

Amendment to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

1. **(Currently Amended)** A membrane translocation peptide carrier moiety consisting of

(a) RRMKWKK (SEQ ID NO: 2);

1 7

(b) SEQ ID No 2, wherein one to three amino acid residues are replaced by a naturally or non-naturally occurring amino acid residue;

(c) SEQ ID No 2, wherein the order of two or more amino acid residues is reversed;

(d) SEQ ID No 2, wherein both (b) and (c) are present together;

(e) SEQ ID No 2, wherein one or more amino acid residues are in peptoid form;

(f) SEQ ID No 2, wherein the (N-C-C) backbone of one or more amino acid residues of the peptide carrier moiety has been modified; or

(g) SEQ ID NO:2, having any of (b)-(f) in combination.

Claims 2-48 (Canceled)

49. **(Currently Amended)** A carrier moiety according to claims 1, 77, 79 or 81, wherein one to three amino acid residues are replaced by homologous replacement.

50. **(Canceled)**

51. **(Currently Amended)** A carrier moiety according to claims 1, 77, 79

or 81, wherein one to three amino acid residues are replaced by non-homologous replacement.

52. **(Canceled)**

53. **(Previously Presented)** A carrier moiety according to claim 51, wherein the replacement amino acid is a non-natural amino acid selected from the group consisting of: alpha* and alpha-disubstituted* amino acids, N-alkyl amino acids*, lactic acid*, halide derivatives of natural amino acids, L-allyl-glycine*, β -alanine*, L- α -amino butyric acid*, L- γ -amino butyric acid*, L- α -amino isobutyric acid*, L- ϵ -amino caproic acid#, 7-amino heptanoic acid*, L-methionine sulfone**, L-norleucine*, L-norvaline*, p-nitro-L-phenylalanine*, L-hydroxyproline#, L-thioproline*, and methyl derivatives of phenylalanine (Phe), L-Phe (4-amino) #, L-Tyr (methyl)*, L-Phe (4-isopropyl)*, L-Tic (1,2,3,4-tetrahydroisoquinoline-3-carboxyl acid)*, L-diaminopropionic acid # and L-Phe (4-benzyl)*, wherein the notation * indicates that the derivative is hydrophobic.

54. **(Currently Amended)** A carrier moiety according to claims 1, 78, 79 or 81, wherein the order of the second and third amino acids from the C-terminal end of the peptide is reversed.

Claims 55-58 **(Canceled)**

59. **(Currently Amended)** A carrier moiety according to claim 1 or 81, wherein one or more amino acid[[s]] residues are in peptoid form.

60. **(Currently Amended)** A carrier moiety according to claims 1, 77, 79 or 81, wherein one to three amino acid residues at any of positions 1, 2, 3, 5, 6 or 7 of said formula (SEQ ID No. 2) are replaced by a naturally or non-naturally occurring amino acid.

61. **(Currently Amended)** A carrier moiety according to claims 1, 78, 79

or 81, wherein the order of two amino acid residues at any of positions 2 and 3, 3 and 4, 4 and 5, or 5 and 6 of said formula (SEQ ID No. 2) are reversed.

62. **(Currently Amended)** A carrier moiety according to claim[[s]] 60, wherein the amino acid residue at position 3 or 7 of said formula (SEQ ID No. 2) is replaced.

63. **(Previously Presented)** A carrier moiety according to claim 60, wherein the amino acid residue at position 3 of said formula (SEQ ID No. 2) is replaced.

64. **(Previously Presented)** A carrier moiety according to claim 61, wherein the order of the amino acid residue at position 3 of said formula (SEQ ID No. 2) is reversed with the amino acid at position 2.

65. **(Previously Presented)** A carrier moiety according to claim 61, wherein the order of the amino acid residue at position 3 of said formula (SEQ ID No. 2) is reversed with the amino acid at position 4.

66. **(Currently Amended)** A carrier moiety according to claim[[s]] 49 wherein homologous replacement occurs at any of positions 1 and 2 of said formula (SEQ ID No. 2).

67. **(Currently Amended)** A carrier moiety according to claim[[s]] 51 [[or 53]], wherein non-homologous replacement occurs at any of positions 3, 4, 5 and 6 of said formula (SEQ ID No. 2).

68. **(Currently Amended)** A carrier moiety according to claims 1, 77, 79 or 81~~claims 1, 49 or 51~~, wherein two amino acid residues of said formula (SEQ ID No. 2) are replaced by homologous or non-homologous replacement.

69. **(Previously Presented)** A carrier moiety according to claim 68,

wherein amino acid residues at positions 2 and 3 of said formula (SEQ ID No. 2) are replaced.

70. **(Previously Presented)** A carrier moiety according to claim 68, wherein amino acid residues at positions 4 and 5 of said formula (SEQ ID No. 2) are replaced.

71. **(Currently Amended)** A carrier moiety according to claim 68, wherein amino acid residues at positions 5 and 6 of said formula (SEQ ID No. 2) are replaced.

72. **(Previously Presented)** A carrier moiety according to claim 53, wherein the halide derivative is selected from the group consisting of trifluorotyrosine*, p-Cl-phenylalanine*, p-Br-phenylalanine*, and p-I-phenylalanine*.

73. **(Previously Presented)** A carrier moiety according to claim 53, wherein the methyl derivative of phenylalanine (Phe) is selected from the group consisting of 4-methyl-Phe*, and pentamethyl-Phe*.

74. **(Currently Amended)** A carrier moiety [[of]]according to claims 1, 76, 77, 78, 79, 80 or 81, wherein the free carboxyl group of the carboxy terminal amino acid residue is in the form -C(O)-NRR', wherein R and R' are each independently selected from the group consisting of: hydrogen, C1-6 alkyl, C1-6 alkylene or C1-6 alkynyl, aryl, each optionally substituted with a heteroatom.

75. **(Currently Amended)** A carrier moiety according to claim 74, wherein the free carboxyl group of the carboxy terminal amino acid residue is a carboxamide group.

76. **(New)** The membrane translocation peptide carrier moiety according to claim 1 consisting of RRMKWKK (SEQ ID NO: 2).

77. **(New)** The membrane translocation peptide carrier moiety according to claim 1, wherein one to three amino acid residues are replaced by a naturally or non-naturally occurring amino acid residue.

78. **(New)** The membrane translocation peptide carrier moiety according to claim 1, wherein the order of two or more amino acid residues is reversed.

79. **(New)** The membrane translocation peptide carrier moiety according to claim 1, wherein one to three amino acid residues are replaced by a naturally or non-naturally occurring amino acid residue and wherein the order of two or more amino acid residues is reversed.

80. **(New)** The membrane translocation peptide carrier moiety according to claim 1, wherein the (N-C-C) backbone of one or more amino acid residues of the peptide carrier moiety has been modified.

81. **(New)** The membrane translocation peptide carrier moiety according to claim 1, having at least two of the following in combination:

(a) SEQ ID No 2, wherein one to three amino acid residues are replaced by a naturally or non-naturally occurring amino acid residue;

(b) SEQ ID No 2, wherein the order of two or more amino acid residues is reversed;

(c) SEQ ID No 2, wherein both (b) and (c) are present together;

(d) SEQ ID No 2, wherein one or more amino acid residues are in peptoid form; or

(e) SEQ ID No 2, wherein the (N-C-C) backbone of one or more amino acid residues of the peptide carrier moiety has been modified.